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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/027,400

DATE: 05/08/2002
TIME: 09:15:21

Input Set : N:\Crf3\RULE60\10027400.raw
Output Set: N:\CRF3\05082002\J027400.raw

```
     SEQUENCE LISTING
 3 (1) GENERAL INFORMATION:
 5      (i) APPLICANT: WILLIAMS, Lewis T.
 6                    ESCOBEDO, Jaime A.
 8     (ii) TITLE OF INVENTION: PLATELET-DERIVED GROWTH FACTOR RECEPTORS
10    (iii) NUMBER OF SEQUENCES: 14
12     (iv) CORRESPONDENCE ADDRESS:
13          (A) ADDRESSEE: Townsend and Townsend and Crew
14          (B) STREET: One Market, Steuart Street Tower, 20th Floor
15          (C) CITY: San Francisco
16          (D) STATE: California
17          (E) COUNTRY: USA
18          (F) ZIP: 94105
20      (v) COMPUTER READABLE FORM:
21          (A) MEDIUM TYPE: Floppy disk
22          (B) COMPUTER: IBM PC compatible
23          (C) OPERATING SYSTEM: PC-DOS/MS-DOS
24          (D) SOFTWARE: PatentIn Release #1.0, Version #1.30
26     (vi) CURRENT APPLICATION DATA:
C-->27          (A) APPLICATION NUMBER: US/10/027,400
C-->28          (B) FILING DATE: 19-Dec-2001
29          (C) CLASSIFICATION:
35    (vii) PRIOR APPLICATION DATA:
32          (A) APPLICATION NUMBER: US/08/461,917
33          (B) FILING DATE: 05-JUN-1995
36          (A) APPLICATION NUMBER: US 07/151,414
37          (B) FILING DATE: 02-FEB-1988
39   (viii) ATTORNEY/AGENT INFORMATION:
40          (A) NAME: Dow, Karen B.
41          (B) REGISTRATION NUMBER: 29,684
42          (C) REFERENCE/DOCKET NUMBER: 2307K-267-2-4
44     (ix) TELECOMMUNICATION INFORMATION:
45          (A) TELEPHONE: 415/326-2400
46          (B) TELEFAX: 415/326-2422
49 (2) INFORMATION FOR SEQ ID NO: 1:
51      (i) SEQUENCE CHARACTERISTICS:
52          (A) LENGTH: 6373 base pairs
53          (B) TYPE: nucleic acid
54          (C) STRANDEDNESS: single
55          (D) TOPOLOGY: linear
57     (ii) MOLECULE TYPE: cDNA
60     (ix) FEATURE:
61          (A) NAME/KEY: CDS
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ENTERED

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62          (B) LOCATION: 129..3398
65     (xi) SEQUENCE DESCRIPTION: SEQ ID NO: 1:
67 TTGGAGCTAC AGGGAGAGAA ACAGAGGAGG AGACTGCAAG AGATCATTGG AGGCCGTGGG     60
69 CACGCTCTTT ACTCCATGTG TGGGACATTC ATTGCGGAAT AACATCGGAG GAGAAGTTTC    120
71 CCAGAGCT ATG GGG ACT TCC CAT CCG GCG TTC CTG GTC TTA GGC TGT CTT    170
72          Met Gly Thr Ser His Pro Ala Phe Leu Val Leu Gly Cys Leu
73            1               5                  10
75 CTC ACA GGG CTG AGC CTA ATC CTC TGC CAG CTT TCA TTA CCC TCT ATC    218
76 Leu Thr Gly Leu Ser Leu Ile Leu Cys Gln Leu Ser Leu Pro Ser Ile
77 15                  20                  25                  30
79 CTT CCA AAT GAA AAT GAA AAG GTT GTG CAG CTG AAT TCA TCC TTT TCT    266
80 Leu Pro Asn Glu Asn Glu Lys Val Val Gln Leu Asn Ser Ser Phe Ser
81                  35                  40                  45
83 CTG AGA TGC TTT GGG GAG AGT GAA GTG AGC TGG CAG TAC CCC ATG TCT    314
84 Leu Arg Cys Phe Gly Glu Ser Glu Val Ser Trp Gln Tyr Pro Met Ser
85              50                  55                  60
87 GAA GAA GAG AGC TCC GAT GTG GAA ATC AGA AAT GAA GAA AAC AAC AGC    362
88 Glu Glu Glu Ser Ser Asp Val Glu Ile Arg Asn Glu Glu Asn Asn Ser
89          65                  70                  75
91 GGC CTT TTT GTG ACG GTC TTG GAA GTG AGC AGT GCC TCG GCG GCC CAC    410
92 Gly Leu Phe Val Thr Val Leu Glu Val Ser Ser Ala Ser Ala Ala His
93      80                  85                  90
95 ACA GGG TTG TAC ACT TGC TAT TAC AAC CAC ACT CAG ACA GAA GAG AAT    458
96 Thr Gly Leu Tyr Thr Cys Tyr Tyr Asn His Thr Gln Thr Glu Glu Asn
97 95                  100                 105                 110
99 GAG CTT GAA GGC AGG CAC ATT TAC ATC TAT GTG CCA GAC CCA GAT GTA    506
100 Glu Leu Glu Gly Arg His Ile Tyr Ile Tyr Val Pro Asp Pro Asp Val
101                 115                 120                 125
103 GCC TTT GTA CCT CTA GGA ATG ACG GAT TAT TTA GTC ATC GTG GAG GAT    554
104 Ala Phe Val Pro Leu Gly Met Thr Asp Tyr Leu Val Ile Val Glu Asp
105                 130                 135                 140
107 GAT GAT TCT GCC ATT ATA CCT TGT CGC ACA ACT GAT CCC GAG ACT CCT    602
108 Asp Asp Ser Ala Ile Ile Pro Cys Arg Thr Thr Asp Pro Glu Thr Pro
109             145                 150                 155
111 GTA ACC TTA CAC AAC AGT GAG GGG GTG GTA CCT GCC TCC TAC GAC AGC    650
112 Val Thr Leu His Asn Ser Glu Gly Val Val Pro Ala Ser Tyr Asp Ser
113     160                 165                 170
115 AGA CAG GGC TTT AAT GGG ACC TTC ACT GTA GGG CCC TAT ATC TGT GAG    698
116 Arg Gln Gly Phe Asn Gly Thr Phe Thr Val Gly Pro Tyr Ile Cys Glu
117 175                 180                 185                 190
119 GCC ACC GTC AAA GGA AAG AAG TTC CAG ACC ATC CCA TTT AAT GTT TAT    746
120 Ala Thr Val Lys Gly Lys Lys Phe Gln Thr Ile Pro Phe Asn Val Tyr
121                 195                 200                 205
123 GCT TTA AAA GCA ACA TCA GAG CTG GAT CTA GAA ATG GAA GCT CTT AAA    794
124 Ala Leu Lys Ala Thr Ser Glu Leu Asp Leu Glu Met Glu Ala Leu Lys
125                 210                 215                 220
127 ACC GTG TAT AAG TCA GGG GAA ACG ATT GTG GTC ACC TGT GCT GTT TTT    842
128 Thr Val Tyr Lys Ser Gly Glu Thr Ile Val Val Thr Cys Ala Val Phe
129                 225                 230                 235
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131 AAC AAT GAG GTG GTT GAC CTT CAA TGG ACT TAC CCT GGA GAA GTG AAA        890
132 Asn Asn Glu Val Val Asp Leu Gln Trp Thr Tyr Pro Gly Glu Val Lys
133     240                 245                 250
135 GGC AAA GGC ATC ACA ATG CTG GAA GAA ATC AAA GTC CCA TCC ATC AAA        938
136 Gly Lys Gly Ile Thr Met Leu Glu Glu Ile Lys Val Pro Ser Ile Lys
137 255                 260                 265                 270
139 TTG GTG TAC ACT TTG ACG GTC CCC GAG GCC ACG GTG AAA GAC AGT GGA        986
140 Leu Val Tyr Thr Leu Thr Val Pro Glu Ala Thr Val Lys Asp Ser Gly
141                 275                 280                 285
143 GAT TAC GAA TGT GCT GCC CGC CAG GCT ACC AGG GAG GTC AAA GAA ATG       1034
144 Asp Tyr Glu Cys Ala Ala Arg Gln Ala Thr Arg Glu Val Lys Glu Met
145                 290                 295                 300
147 AAG AAA GTC ACT ATT TCT GTC CAT GAG AAA GGT TTC ATT GAA ATC AAA       1082
148 Lys Lys Val Thr Ile Ser Val His Glu Lys Gly Phe Ile Glu Ile Lys
149     305                 310                 315
151 CCC ACC TTC AGC CAG TTG GAA GCT GTC AAC CTG CAT GAA GTC AAA CAT       1130
152 Pro Thr Phe Ser Gln Leu Glu Ala Val Asn Leu His Glu Val Lys His
153     320                 325                 330
155 TTT GTT GTA GAG GTG CGG GCC TAC CCA CCT CCC AGG ATA TCC TGG CTG       1178
156 Phe Val Val Glu Val Arg Ala Tyr Pro Pro Pro Arg Ile Ser Trp Leu
157 335                 340                 345                 350
159 AAA AAC AAT CTG ACT CTG ATT GAA AAT CTC ACT GAG ATC ACC ACT GAT       1226
160 Lys Asn Asn Leu Thr Leu Ile Glu Asn Leu Thr Glu Ile Thr Thr Asp
161                 355                 360                 365
163 GTG GAA AAG ATT CAG GAA ATA AGG TAT CGA AGC AAA TTA AAG CTG ATC       1274
164 Val Glu Lys Ile Gln Glu Ile Arg Tyr Arg Ser Lys Leu Lys Leu Ile
165                 370                 375                 380
167 CGT GCT AAG GAA GAA GAC AGT GGC CAT TAT ACT ATT GTA GCT CAA AAT       1322
168 Arg Ala Lys Glu Glu Asp Ser Gly His Tyr Thr Ile Val Ala Gln Asn
169             385                 390                 395
171 GAA GAT GCT GTG AAG AGC TAT ACT TTT GAA CTG TTA ACT CAA GTT CCT       1370
172 Glu Asp Ala Val Lys Ser Tyr Thr Phe Glu Leu Leu Thr Gln Val Pro
173     400                 405                 410
175 TCA TCC ATT CTG GAC TTG GTC GAT GAT CAC CAT GGC TCA ACT GGG GGA       1418
176 Ser Ser Ile Leu Asp Leu Val Asp Asp His His Gly Ser Thr Gly Gly
177 415                 420                 425                 430
179 CAG ACG GTG AGG TGC ACA GCT GAA GGC ACG CCG CTT CCT GAT ATT GAG       1466
180 Gln Thr Val Arg Cys Thr Ala Glu Gly Thr Pro Leu Pro Asp Ile Glu
181                 435                 440                 445
183 TGG ATG ATA TGC AAA GAT ATT AAG AAA TGT AAT AAT GAA ACT TCC TGG       1514
184 Trp Met Ile Cys Lys Asp Ile Lys Lys Cys Asn Asn Glu Thr Ser Trp
185                 450                 455                 460
187 ACT ATT TTG GCC AAC AAT GTC TCA AAC ATC ATC ACG GAG ATC CAC TCC       1562
188 Thr Ile Leu Ala Asn Asn Val Ser Asn Ile Ile Thr Glu Ile His Ser
189             465                 470                 475
191 CGA GAC AGG AGT ACC GTG GAG GGC CGT GTG ACT TTC GCC AAA GTG GAG       1610
192 Arg Asp Arg Ser Thr Val Glu Gly Arg Val Thr Phe Ala Lys Val Glu
193     480                 485                 490
195 GAG ACC ATC GCC GTG CGA TGC CTG GCT AAG AAT CTC CTT GGA GCT GAG       1658
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                    RAW SEQUENCE LISTING              DATE: 05/08/2002
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196 Glu Thr Ile Ala Val Arg Cys Leu Ala Lys Asn Leu Leu Gly Ala Glu
197 495                 500             505                 510
199 AAC CGA GAG CTG AAG CTG GTG GCT CCC ACC CTG CGT TCT GAA CTC ACG      1706
200 Asn Arg Glu Leu Lys Leu Val Ala Pro Thr Leu Arg Ser Glu Leu Thr
201                 515             520                 525
203 GTG GCT GCT GCA GTC CTG GTG CTG TTG GTG ATT GTG ATC ATC TCA CTT      1754
204 Val Ala Ala Ala Val Leu Val Leu Leu Val Ile Val Ile Ile Ser Leu
205                 530             535                 540
207 ATT GTC CTG GTT GTC ATT TGG AAA CAG AAA CCG AGG TAT GAA ATT CGC      1802
208 Ile Val Leu Val Val Ile Trp Lys Gln Lys Pro Arg Tyr Glu Ile Arg
209             545             550                 555
211 TGG AGG GTC ATT GAA TCA ATC AGC CCA GAT GGA CAT GAA TAT ATT TAT      1850
212 Trp Arg Val Ile Glu Ser Ile Ser Pro Asp Gly His Glu Tyr Ile Tyr
213         560             565             570
215 GTG GAC CCG ATG CAG CTG CCT TAT GAC TCA AGA TGG GAG TTT CCA AGA      1898
216 Val Asp Pro Met Gln Leu Pro Tyr Asp Ser Arg Trp Glu Phe Pro Arg
217 575                 580             585                 590
219 GAT GGA CTA GTG CTT GGT CGG GTC TTG GGG TCT GGA GCG TTT GGG AAG      1946
220 Asp Gly Leu Val Leu Gly Arg Val Leu Gly Ser Gly Ala Phe Gly Lys
221             595             600                 605
223 GTG GTT GAA GGA ACA GCC TAT GGA TTA AGC CGG TCC CAA CCT GTC ATG      1994
224 Val Val Glu Gly Thr Ala Tyr Gly Leu Ser Arg Ser Gln Pro Val Met
225             610             615                 620
227 AAA GTT GCA GTG AAG ATG CTA AAA CCC ACG GCC AGA TCC AGT GAA AAA      2042
228 Lys Val Ala Val Lys Met Leu Lys Pro Thr Ala Arg Ser Ser Glu Lys
229             625             630                 635
231 CAA GCT CTC ATG TCT GAA CTG AAG ATA ATG ACT CAC CTG GGG CCA CAT      2090
232 Gln Ala Leu Met Ser Glu Leu Lys Ile Met Thr His Leu Gly Pro His
233         640             645             650
235 TTG AAC ATT GTA AAC TTG CTG GGA GCC TGC ACC AAG TCA GGC CCC ATT      2138
236 Leu Asn Ile Val Asn Leu Leu Gly Ala Cys Thr Lys Ser Gly Pro Ile
237 655             660             665                 670
239 TAC ATC ATC ACA GAG TAT TGC TTC TAT GGA GAT TTG GTC AAC TAT TTG      2186
240 Tyr Ile Ile Thr Glu Tyr Cys Phe Tyr Gly Asp Leu Val Asn Tyr Leu
241             675             680                 685
243 CAT AAG AAT AGG GAT AGC TTC CTG AGC CAC CAC CCA GAG AAG CCA AAG      2234
244 His Lys Asn Arg Asp Ser Phe Leu Ser His His Pro Glu Lys Pro Lys
245             690             695             700
247 AAA GAG CTG GAT ATC TTT GGA TTG AAC CCT GCT GAT GAA AGC ACA CGG      2282
248 Lys Glu Leu Asp Ile Phe Gly Leu Asn Pro Ala Asp Glu Ser Thr Arg
249         705             710             715
251 AGC TAT GTT ATT TTA TCT TTT GAA AAC AAT GGT GAC TAC ATG GAC ATG      2330
252 Ser Tyr Val Ile Leu Ser Phe Glu Asn Asn Gly Asp Tyr Met Asp Met
253         720             725             730
255 AAG CAG GCT GAT ACT ACA CAG TAT GTC CCC ATG CTA GAA AGG AAA GAG      2378
256 Lys Gln Ala Asp Thr Thr Gln Tyr Val Pro Met Leu Glu Arg Lys Glu
257 735             740             745                 750
259 GTT TCT AAA TAT TCC GAC ATC CAG AGA TCA CTC TAT GAT CGT CCA GCC      2426
260 Val Ser Lys Tyr Ser Asp Ile Gln Arg Ser Leu Tyr Asp Arg Pro Ala
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261             755              760              765
263 TCA TAT AAG AAG AAA TCT ATG TTA GAC TCA GAA GTC AAA AAC CTC CTT      2474
264 Ser Tyr Lys Lys Lys Ser Met Leu Asp Ser Glu Val Lys Asn Leu Leu
265             770              775              780
267 TCA GAT GAT AAC TCA GAA GGC CTT ACT TTA TTG GAT TTG TTG AGC TTC      2522
268 Ser Asp Asp Asn Ser Glu Gly Leu Thr Leu Leu Asp Leu Leu Ser Phe
269             785              790              795
271 ACC TAT CAA GTT GCC CGA GGA ATG GAG TTT TTG GCT TCA AAA AAT TGT      2570
272 Thr Tyr Gln Val Ala Arg Gly Met Glu Phe Leu Ala Ser Lys Asn Cys
273             800              805              810
275 GTC CAC CGT GAT CTG GCT GCT CGC AAC GTT CTC CTG GCA CAA GGA AAA      2618
276 Val His Arg Asp Leu Ala Ala Arg Asn Val Leu Leu Ala Gln Gly Lys
277 815             820              825              830
279 ATT GTG AAG ATC TGT GAC TTT GGC CTG GCC AGA GAC ATC ATG CAT GAT      2666
280 Ile Val Lys Ile Cys Asp Phe Gly Leu Ala Arg Asp Ile Met His Asp
281             835              840              845
283 TCG AAC TAT GTG TCG AAA GGC AGT ACC TTT CTG CCC GTG AAG TGG ATG      2714
284 Ser Asn Tyr Val Ser Lys Gly Ser Thr Phe Leu Pro Val Lys Trp Met
285             850              855              860
287 GCT CCT GAG AGC ATC TTT GAC AAC CTC TAC ACC ACA CTG AGT GAT GTC      2762
288 Ala Pro Glu Ser Ile Phe Asp Asn Leu Tyr Thr Thr Leu Ser Asp Val
289             865              870              875
291 TGG TCT TAT GGC ATT CTG CTC TGG GAG ATC TTT TCC CTT GGT GGC ACC      2810
292 Trp Ser Tyr Gly Ile Leu Leu Trp Glu Ile Phe Ser Leu Gly Gly Thr
293 880             885              890
295 CCT TAC CCC GGC ATG ATG GTG GAT TCT ACT TTC TAC AAT AAG ATC AAG      2858
296 Pro Tyr Pro Gly Met Met Val Asp Ser Thr Phe Tyr Asn Lys Ile Lys
297 895             900              905              910
299 AGT GGG TAC CGG ATG GCC AAG CCT GAC CAC GCT ACC AGT GAA GTC TAC      2906
300 Ser Gly Tyr Arg Met Ala Lys Pro Asp His Ala Thr Ser Glu Val Tyr
301             915              920              925
303 GAG ATC ATG GTG AAA TGC TGG AAC AGT GAG CCG GAG AAG AGA CCC TCC      2954
304 Glu Ile Met Val Lys Cys Trp Asn Ser Glu Pro Glu Lys Arg Pro Ser
305             930              935              940
307 TTT TAC CAC CTG AGT GAG ATT GTG GAG AAT CTG CTG CCT GGA CAA TAT      3002
308 Phe Tyr His Leu Ser Glu Ile Val Glu Asn Leu Leu Pro Gly Gln Tyr
309             945              950              955
311 AAA AAG AGT TAT GAA AAA ATT CAC CTG GAC TTC CTG AAG AGT GAC CAT      3050
312 Lys Lys Ser Tyr Glu Lys Ile His Leu Asp Phe Leu Lys Ser Asp His
313 960             965              970
315 CCT GCT GTG GCA CGC ATG CGT GTG GAC TCA GAC AAT GCA TAC ATT GGT      3098
316 Pro Ala Val Ala Arg Met Arg Val Asp Ser Asp Asn Ala Tyr Ile Gly
317 975             980              985              990
319 GTC ACC TAC AAA AAC GAG GAA GAC AAG CTG AAG GAC TGG GAG GGT GGT      3146
320 Val Thr Tyr Lys Asn Glu Glu Asp Lys Leu Lys Asp Trp Glu Gly Gly
321             995              1000             1005
323 CTG GAT GAG CAG AGA CTG AGC GCT GAC AGT GGC TAC ATC ATT CCT CTG      3194
324 Leu Asp Glu Gln Arg Leu Ser Ala Asp Ser Gly Tyr Ile Ile Pro Leu
325             1010             1015             1020
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VERIFICATION SUMMARY   DATE: 05/08/2002
PATENT APPLICATION: US/10/027,400   TIME: 09:15:22

Input Set : **N:\Crf3\RULE60\10027400.raw**
Output Set: **N:\CRF3\05082002\J027400.raw**

L:870 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:874 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:878 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:882 M:336 W: Invalid Amino Acid Number in Coding Region, SEQ ID:3
L:1493 M:246 W: Invalid value of Alpha Sequence Header Field, [MOLECULE TYPE:], SeqNo=14

A DOCPHOENIX

APPL PARTS

_____ IMIS _____
Internal Misc. Paper

_____ LET. _____
Misc. Incoming Letter

_____ 371P _____
PCT Papers in a 371Application

_____ A... _____
Amendment Including Elections

_____ ABST _____
Abstract

_____ ADS _____
Application Data Sheet

_____ AF/D _____
Affidavit or Exhibit Received

_____ APPENDIX _____
Appendix

_____ ARTIFACT _____
Artifact

_____ BIB _____
Bib Data Sheet

_____ CLM _____
Claim

_____ COMPUTER _____
Computer Program Listing

_____ CRFL _____
All CRF Papers for Backfile

_____ DIST _____
Terminal Disclaimer Filed

_____ DRW _____
Drawings

_____ FOR _____
Foreign Reference

_____ FRPR _____
Foreign Priority Papers

_____ IDS _____
IDS Including 1449

_____ NPL _____
Non-Patent Literature

_____ OATH _____
Oath or Declaration

_____ PET. _____
Petition

_____ RETMAIL _____
Mail Returned by USPS

_____ SEQLIST _____
Sequence Listing

_____ SPEC _____
Specification

_____ SPEC NO _____
Specification Not in English

_____ TRNA _____
Transmittal New Application

OUTGOING

_____ CTMS _____
Misc. Office Action

_____ 1449 _____
Signed 1449

_____ 892 _____
892

_____ ABN _____
Abandonment

_____ APDEC _____
Board of Appeals Decision

_____ APEA _____
Examiner Answer

_____ CTAV _____
Count Advisory Action

_____ CTEQ _____
Count Ex parte Quayle

_____ CTFR _____
Count Final Rejection

_____ CTNF _____
Count Non-Final

_____ CTRS _____
Count Restriction

_____ EXIN _____
Examiner Interview

_____ M903 _____
DO/EO Acceptance

_____ M905 _____
DO/EO Missing Requirement

_____ NFDR _____
Formal Drawing Required

_____ NOA _____
Notice of Allowance

_____ PETDEC _____
Petition Decision

INCOMING

_____ AP.B _____
Appeal Brief

_____ C.AD _____
Change of Address

_____ N/AP _____
Notice of Appeal

_____ PA.. _____
Change in Power of Attorney

_____ REM _____
Applicant Remarks in Amendment

_____ XT/ _____
Extension of Time filed separate

Internal

_____ SRNT _____
Examiner Search Notes

_____ CLMPTO _____
PTO Prepared Complete Claim Set

_____ ECBOX _____
Evidence Copy Box Identification

_____ WCLM _____
Claim Worksheet

_____ WFEE _____
Fee Worksheet

File Wrapper

_____ FWCLM _____
File Wrapper Claim

_____ IIFW _____
File Wrapper Issue Information

_____ SRFW _____
File Wrapper Search Info